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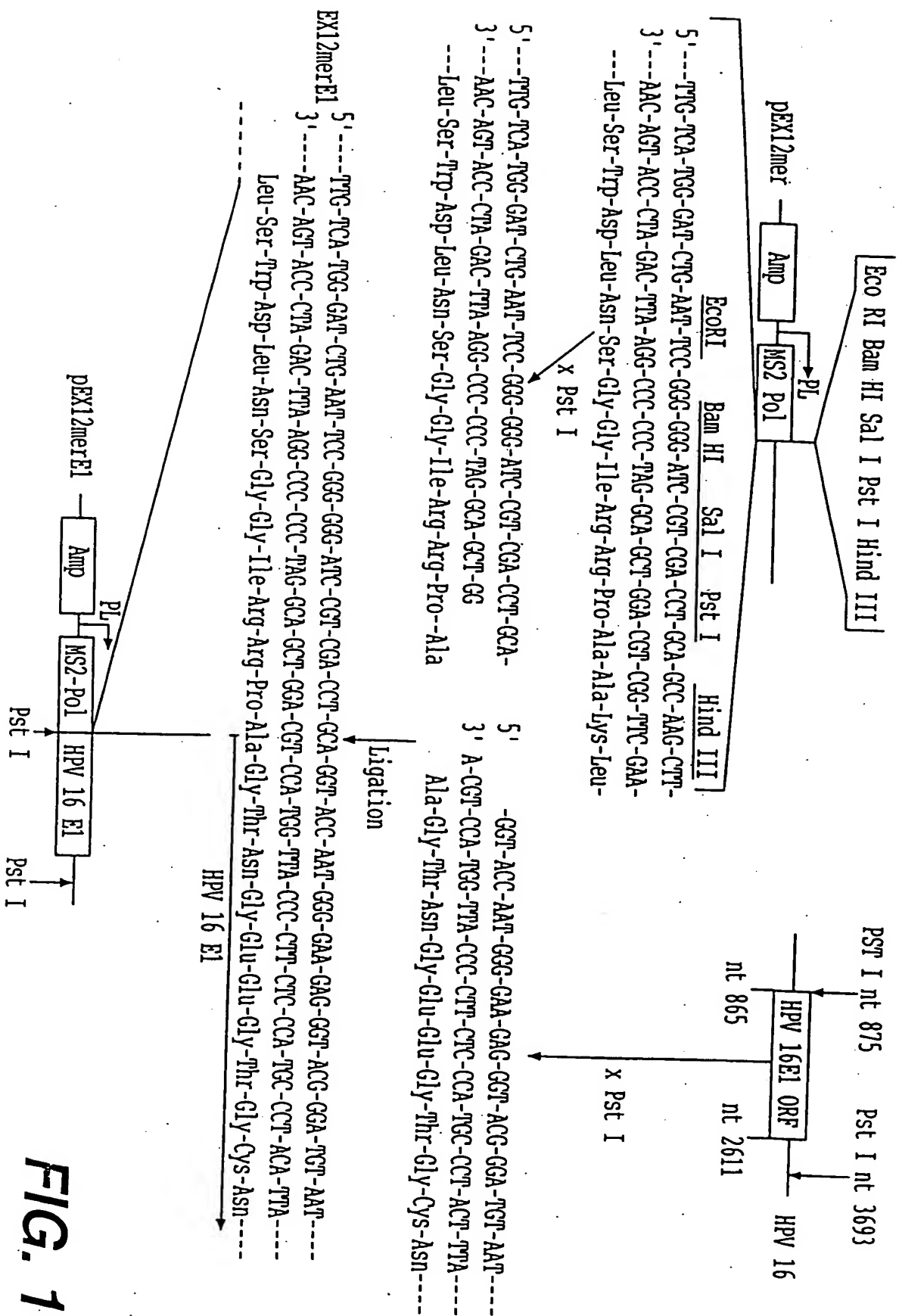
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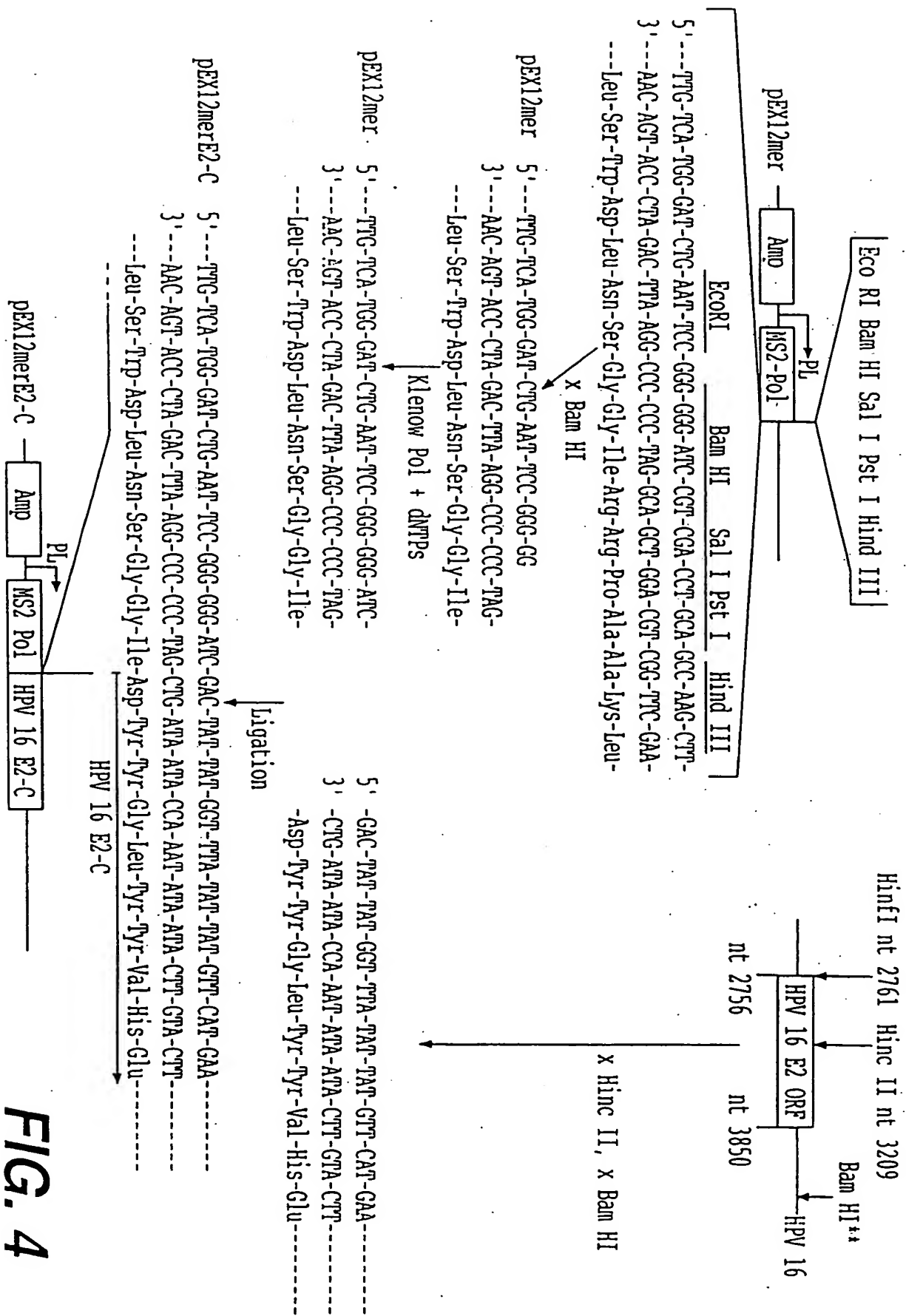
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CLONE	SEQUENCE	POSITION	ISOLATE
<u>REGION E1-1090:</u>			
1090	NGMFTVEAVEKKTGDAISDDENENDSPGCDLVDFIWNNDYLT	AS 16-60	1
1079	NENDSPGCDLVDFIWN	AS 38-55	3
1084	HADPAGTNGEEGTCNGMFTVEAVEVEKKTGDAISDDENENDSPGCDLVDFIWNNDYLT	AS 1-60	3
1029	EDLVDFIWNNDYLT	AS 46-60	5
1099	EDLVDFIWNNDYLTQAETETAAHALFTAQEAQH	AS 46-79	5
1145	NENDSPGCDLVDFIWNNDYLTQAETETAAHALFTAQEAQHDAVQVLKRYLWVh	AS 38-91	1
<u>REGION E1-1059:</u>			
1059	stgsktkvfGSPPLSDIS	AS 92-99	1

FIG. 2





**FIG. 4**

CLONE	SEQUENCE	POSITION	ISOLATE
<u>REGION E2-1066:</u>			
1066	DKILTHYENDS	AS 13-23	3
1025	DKILTHYENDSTDLRDHI	AS 13-30	5
<u>REGION E2-1170:</u>			
1170	DLRHDIDYKKA	AS 25-35	2
<u>REGION E2-1074:</u>			
1074	AIYYKARENGCFKHINHQPPTLA	AS 41-63	4
1060	AIYYKARENGCFKHINHQPPTLAVSKKAL	AS 41-70	2
1061	YYKARENGCFKHINHQPPTLAVSKN	AS 43-67	1
1063	INHQPPTLAVSKKALQAI	AS 55-73	1
1057	INHQPPTLAVSKKAL	AS 55-70	4
1018	TLAVSKKALQAIQLTLETINSQYSNEKMTLQDV	AS 61-76	2
<u>REGION E2-1112:</u>			
1112	QLTLETINSQYSNEKMTLQDVSL	AS 76-100	1
1156	TLTETINSQYSNEK	AS 78-91	1
<u>REGION E2-1158:</u>			
1158	TSVFSSNEVSSPEII	AS 197-211	1
1121	VFSSNEVSSPEIIRQHLANHPAHTPKAVALGTEET	AS 199-234	2
<u>REGION E2-1102:</u>			
1102	EIIRQHLANHPAHTPKAVALGTEETQTTIORPRSEP	AS 209-245	1
1091	TEETQTTIORPRSEPDCN	AS 231-249	2

FIG. 5